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(FILE 'HOME' ENTERED AT 13:52:27 ON 30 JUL 2007)

FILE 'MEDLINE' ENTERED AT 13:53:26 ON 30 JUL 2007

L1 55999 S SEQUENCE ALIGNMENT
L2 100985 S ALGORITHM
L3 15 S SIMULTANEOUSLY MUTATED
L4 669 S PHYLOGENETICS
L5 0 S (SIMULTANEOUSLY MUTATED) AND (SEQUENCE ALIGNMENT)
L6 14 S (SEQUENCE ALIGNMENT) AND ALGORITHM AND PHYLOGENETICS
L7 0 S (SEQUENCE ALIGNMENT) AND ALGORITHM AND PHYLOGENETICS AND (HIV
L8 10 S (SEQUENCE ALIGNMENT) AND ALGORITHM AND (HIV PROTEASE)
L9 33785 S MUTATIONAL ANALYSIS
L10 382461 S RESISTANCE
L11 1235 S (MUTATIONAL ANALYSIS) AND RESISTANCE
L12 7940 S HIV PROTEASE
L13 21 S (MUTATIONAL ANALYSIS) AND RESISTANCE AND (HIV PROTEASE)
L14 3 S (MUTATIONAL ANALYSIS) AND RESISTANCE AND (HIV PROTEASE) AND (
L15 1660 S HYPERMUTATION
L16 4 S HYPERMUTATION AND (HIV PROTEASE)
L17 14 S HYPERMUTATION AND ALGORITHM
L18 1 S HYPERMUTATION AND (HIV PROTEASE) AND (SEQUENCE ALIGNMENT)
L19 30 S RESISTANCE AND (HIV PROTEASE) AND (SEQUENCE ALIGNMENT)
L20 3 S RESISTANCE AND (HIV PROTEASE) AND (SEQUENCE ALIGNMENT) AND MU